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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)
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Reviewer: Keisha Douglas

Timestamp: [year=2008; month=2; day=28; hr=18; min=26; sec=55; ms=252;]

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Reviewer Comments:

<210> 5

<211> 16

<212> PRT

<213> Artificial

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<223> characteristic sequence motif construct

The above <223> response for sequence id# 5 is invalid, please explain
source of genetic material.

Application No: 10555587 Version No: 2.0

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Output Set:

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Finished: 2008-02-14 17:33:23.494
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 633 ms
Total Warnings: 5
Total Errors: 0
No. of SeqIDs Defined: 5
Actual SeqID Count: 5

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W 213	Artificial or Unknown found in <213> in SEQ ID (5)

SEQUENCE LISTING

<110> Jones, Brian Edward
 Grant, William D.
 Heaphy, Shaun
 Rees, Helen C.
 Grant, Susan

<120> Novel Lipolytic Enzyme ELIP

<130> GC800-2-US

<140> 10555587
 <141> 2008-02-14

<150> PCT/US2004/014685
 <151> 2004-05-12

<150> US 60/470,069
 <151> 2003-05-12

<160> 5

<170> PatentIn version 3.4

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 <223> Xaa can be any naturally occurring amino acid

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 Thr Val Asn Tyr Arg Leu Gly His Leu Gly Phe Phe Ala His Pro Ala
 35 40 45
 Leu Asp Glu Glu Asn Pro Asp Gly Pro Val His Asn Phe Ala Leu Leu
 50 55 60
 Asp Gln Ile Ala Ala Leu Lys Trp Val Gln Glu Asn Ile Ala Ala Phe
 65 70 75 80
 Gly Gly Asp Ala Gly Asn Val Thr Leu Phe Gly Glu Ser Ala Gly Ala
 85 90 95
 Arg Ser Val Leu Ser Leu Leu Ala Ser Pro Leu Ala Lys Asn Leu Phe
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 His Lys Gly Ile Ile Gln Ser Ala Tyr Thr Leu Pro Asp Val Asp Arg
 115 120 125
 Lys Lys Ala Leu Lys Arg Gly Val Ala Leu Ala Gly His Tyr Gly Leu
 130 135 140
 Gln Asn Ala Thr Ala Asp Glu Leu Arg Ala Leu Pro Ala Asp Gly Leu
 145 150 155 160
 Trp Ala Leu Glu Gly Pro Leu Asn Ile Gly Pro Thr Pro Ile Ser Gly
 165 170 175
 Asp Val Val Leu Pro Glu Pro Met Leu Asp Ile Phe Phe Ala Gly Arg
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 Gln His Arg Met Pro Leu Met Val Gly Ser Asn Ser Asp Glu Ala Ser
 195 200 205
 Val Leu Ser Tyr Phe Gly Ile Asp Pro Ala Gly Gln Val Glu Leu Leu
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 Arg Arg Gly Ala Ala Phe Pro Asp Trp Gly Leu Ile Lys Leu Leu Tyr
 225 230 235 240
 Ser Arg Ser Glu Xaa Gly Met Pro Glu Leu Gly Arg Gln Val Cys Arg
 245 250 255
 Asp Met Ala Phe Xaa Xaa Leu Gly Phe Val Val Met Gln Ala Gln Gln
 260 265 270
 Arg Val Asn Gln Pro Cys Trp Arg Tyr Tyr Phe Asp Tyr Val Gly Glu
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 Ala Glu Arg Lys Ile Tyr Ala Asn Gly Thr Trp His Gly Asn Glu Val
 290 295 300
 Pro Tyr Val Phe Asp Thr Leu Ser Leu Thr Pro Pro Ala Ser Glu Tyr
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Val Asn Gln Asn Asp Leu Thr Phe Ala Gly Gln Ile Cys Asp Tyr Trp
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 Thr Arg Phe Ala Arg Ser Ala Gly Pro His Ser Lys Ala Ile Pro Gly
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 Pro Leu Ser Trp Pro Ala Cys Val Arg Gly Lys Asp Arg Thr Met Arg
 355 360 365
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